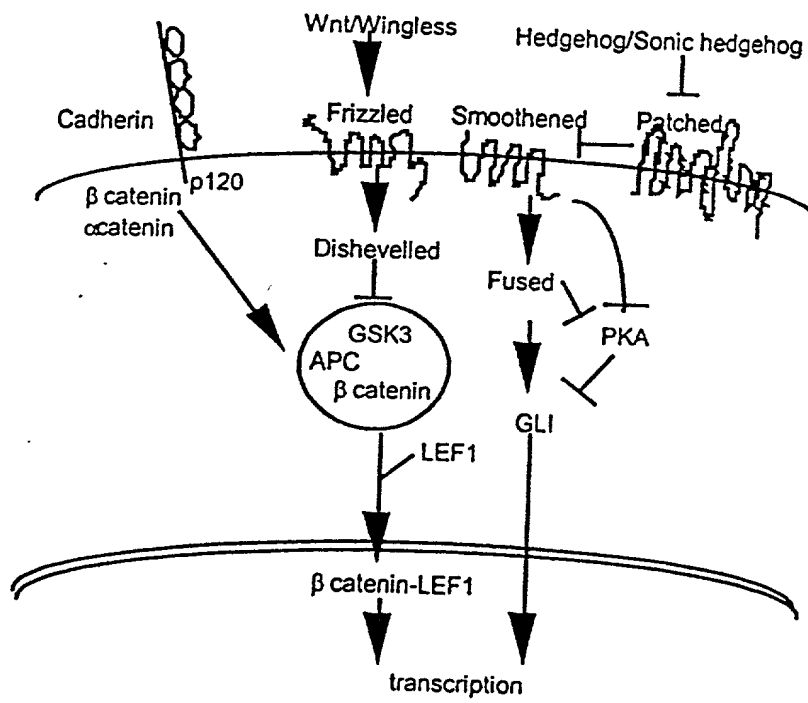


Figure 1



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Figure 2a

Alignment of several frizzled family members

	→	amino terminal domain
fz3/mouse	-----	MAVSWIVFDLWLLTVFLG---QIGGHS-----LFSCE
fz4/mouse	-----	MAWPGTGPS---SRGAPGGVGLRLGLLLQFLLLLRPTLGFGD-----EEERRCD
fz8/mouse	-----	MEWGYL-----LEVTSLLAALAVLQRSSG-AAAASAK-----ELACQ
fz5/human	-----	MARPDP-----SAPPSLL--LLLLLAQLVG-RAAAASK-----APVCQ
fzd9/human	-----	MAVAPLRGALLLWQLLAAGGALEIGRFD-----PERGRG-----AAPCQ
fz1/rat		LEAPLLLGVRAQPAG---QVSG-PGQQRPPPPQPQGG--QQYNGERG--ISIPDHGYCQ
fz2/rat	-----	MRARSAL---PRSALRLLLLPLLLLPAAGP--AQFHGEKG--ISIPDHGFCQ
fz/Dros		ILPTLIQGVQRYDQS---PLDASPYRSGGGLMASSG---TELDG-----LPHHNRCE
fz2/Dros/		GLVLLLTSCRADGPL-----HSADHGMGGMGMGGHGLD-ASPAPGYGVPAPKDPNLRCE
		*:
		CRD
fz3/mouse		PITLRMCQDLPYNTTFMPNLLNHYDQQTAALAMEPFHMPVNLDCSRDFRPFLCALYAPIC
fz4/mouse		PIRIAMCQNLGYNVTKMPNLVGHLELQTDALQLTTFTPLIQYGCSSQLQFFLCSVYVPMC
fz8/mouse		EITVPLCKGIGYNYTYMPNQFNHDTQDEAGLEVHQFWPLVEIQCSDDLKFFLCSMYTPIC
fz5/human		EITVPMCRGIGYNLTHMPNQFNHDTQDEAGLEVHQFWPLVEIQCSDDLKFFLCSMYTPIC
fzd9/human		AVEIPMCRGIGYNLTRMPNLLGHTSQGEAAAEALAEFAPLVQYGCCHSLRFFLCSLYAPMC
fz1/rat		PISIPLCIDIAYNQITIMPNNLLGHTNQEDAGLEVHQFYPLVKVQCSAELKFFLCSMYAPVC
fz2/rat		PISIPLCIDIAYNQITIMPNNLLGHTNQEDAGLEVHQFYPLVKVQCSPELRFLLCSMYAPVC
fz/Dros		PITISICKNIPYNMTIMPNNLIGHTKQEEAGLEVHQFAPLVKIGCSSDDLQFLCSLYVPVC
fz2/Dros/		EITIPMCRGIGYNMTSFPNEMNHETQDEAGLEVHQFWPLVEIKCSDDLKFFLCSMYTPIC
		: : * . : ** : ** . * * * : : * : : * : : ** : **
		CRD
fz3/mouse		M-EYGRVTLPCCRLLCQRAYSECSKLMEMFG-VPWPEDMECSRFPDCLD-EPYPRIVDLN--
fz4/mouse		TEKINIPIGPCGGMCLSVKRRCEPVLREFG-FAWPDTLNCSKFPPQN-DHNHMCMEGP--
fz8/mouse		LEDYKKPLPPCRSVCERAKAGCAPLMRQYG-FAWPDRMRCDLRLEQG-NPDTLCMDYN-R
fz5/human		LPDYHKPLPPCRSVCERAKAGCSPLMRQYG-FAWPERMSCDLRPLVGRDAEVLCMDYN-R
fzd9/human		TDQVSTPIPACRPMCEQARLRCAPIMEQFN-FGWPDSLDCARLPTRN-DPHALCMEAPEN
fz1/rat		T-VLEQALPPCRSLCERA-QGCEALMNKFG-FQWPDTLKCEKFPVHG--AGELCVGQNTS
fz2/rat		T-VLEQAIPPCRSICERARQGCEALMNKFG-FQWPERLRCEHFPRHG--AEQICVGQNHS
fz/Dros		T-ILERPIPPCRSLCESA-RVCEKLMKTYN-FNWPENLECSKFPVHG--GEDLCVAENTT
fz2/Dros/		LEDYHKPLPVCRSVCERARSGCAPIMQOYS-FEWPERMACEHLPLHG-DPDNLCMEQPSY
		* : * . * ** . : * : *
		:
		:
fz3/mouse	-----	LVGDPTTE---
fz4/mouse	---GDEE-----	VPLPHKTP-
fz8/mouse	TDLTAAAPSPRRLLPPPPPPGEQPPSGSGHSRPPGARPPHGGSSRSGDAAAAPPSRGG	
fz5/human	SEATTAPPRP---FPAKP---TLP--G-----PP-----G---APAS-GG	
fzd9/human	ATAGPAEPHK---GLGM---LP-----VAPRPARPPG	
fz1/rat	DKGTPTPSL-----L-----PEFWTSNPQHG	
fz2/rat	EDG--TPAL-----L-----TTAPPSGLQPG	
fz/Dros	SSA-----STAATPTRSVA	
fz2/Dros/	TEAGSGGSSG---GSGG---SGSGSGSGGKRKQGGSGSGSGAGGSSGSTSTKPCR-GR	

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Figure 2b

amino terminal domain continued

		← TM1 →
fz3/mouse	YSFLHVRDCSPPCPN-----MYFRREELSFARYFIGLISIICLSATLFTFLTLFLIDVTR	
fz4/mouse	--LNCVLKCGYDAG-----LYSRSAKEFTDIWMAVWASLCFISTTFTVLTLFLIDSSR	
fz8/mouse	-KTGQIANCALPCHN-----PFFSQDERAFTVFWIGLWSVLCFVSTFATVSTFLIDMER	
fz5/human	-RTGQVPNCAPCYQ-----PSFSADERTFATFWIGLWSVLCFISTSTTVATFLIDMDT	
fzd9/human	--RSCAPRCGPGVEV-----FWSRRDKDFALVWMAVWSALCFFSTAFTVLTLFLEPHR	
fz1/rat	LGEK---DCGAPCEPTKVYGLMYFGPEELRFSRTWIGIWSVLCFASTLFTVLTYLVDMRR	
fz2/rat	LGER---DCAAPCEPARPDGSMFFSHHTRFARLWILTWSVLCFASTFTVTTSVLAMQR	
fz/Dros	VGGKDLHDGAPCH-----AMFFPERERTVLRYWVGSWAAVCVASCLFTVLTLFLIDSSR	
fz2/Dros/	QRIAGVPNCGIPCKG-----PFFSNDEKDFAGLWIALWSGLCFCSTLMTLTTFIIDTER	
	* : : *	

	← TM2 →	extracellular domain loop 1
fz3/mouse	FRYPERPIIFYAVCYMMVSLIFFIGFLE-DRVACNASSP-----	
fz4/mouse	FSYPERPIIFLSMCYNIYSIAYIVRLTVGRERISCDF-----	
fz8/mouse	FKYPERPIIFLSACYLFSVGYLVRVLVAGHEKVACSGGAPGAGGRGGAGGAAAAGAGAAG	
fz5/human	FRYPERPIIFLSACYLCVSLGFLVRLVVGHASVACS-----	
fzd9/human	FQYPERPIIFLSMCYNVYSIAFLIRAVAGAQSACD-----	
fz1/rat	FSYPERPIIFLSGCTAVAVAYIAGFLE-DRVVCNDKFAE-----	
fz2/rat	FRYPERPIIFLSGCTMVSVAIYAGFVLQ-ERVVCNERFSE-----	
fz/Dros	FRYPERAIVFLAVCYLVGCAIVAGLGAG-DSVSCREPFPPPVK---LG-----	
fz2/Dros/	FKYPERPIVFLSACYFMVAVGYLSRNLQNEEIIACDG-----	
	* **** : : *	

	← TM3 →
fz3/mouse	-----AQYKASTVTQGSNKK-ACTMLFMVLYFFTMAGSVWVWVILTITWFLA
fz4/mouse	-----EEAAEPVLIQEGKNTGCAIIIFLLMYFFGMASSIWWVILTLTWFLA
fz8/mouse	RGASSPGARGEYEELGAVEQHVRVYETGPALCTVVFLLVYFFGMASSIWWVILSLTWFLA
fz5/human	R-----EHNHIHYETGPALCTIVFLLVYFFGMASSIWWVILSLTWFLA
fzd9/human	-----QEAGALYVIQEGLENTGCTLVFLLLYFFGMASSLWVWVLTTLTWFLA
fz1/rat	-----DGARTVAQGTKKE-GCTILFMMLYFFSMASSIWWVILSLTWFLA
fz2/rat	-----DGYRTVGQGTKKE-GCTILFMMLYFFSMASSIWWVILSLTWFLA
fz/Dros	-----RLQMMSTITQHRQTTCTVLFMALYFCCMAAFWWSCLAFAWFLA
fz2/Dros/	-----LLLRESSTGPHSCTLVFLLTYFFGMASSIWWVILTITWFLA
	: * : : : *

	← TM4 →	extracellular domain loop 2
fz3/mouse	AVPKWGSEAIKALLFHASAWGIPGTLTIIILLAMNKIEGDNISGVCVGLYDVLDALRYF	
fz4/mouse	AGLKWGHEAIEMHSSYFHIAAWAIPAVKTIVILIMRLVDADELTLGLCYVGNQNLDAITGF	
fz8/mouse	AGMKWGNEAIIAGYSQYFHLAAWLVPVSVKSIIVLALSSVDGDPVAGICYVGNQSLDNLRGF	
fz5/human	AAMKWGNEAIIAGYQYFHLAAWLIPSVKSITALLALSSVDGDPVAGICYVGNQNLNLSLRRF	
fzd9/human	AGKKWGHEAIEAHGSYFHMAAWGLPALKTIVILTIRKVAGDELTLGLCYVASTDAAALTGF	
fz1/rat	AGMKWGHEAIEANSQYFHLAAWVPAIKTITILALGQVDGDLVSGVCVGLNNDALRGF	
fz2/rat	AGMKWGHEAIEANSQYFHLAAWVPAVKTITILAMGQIDGDLVSGVCVGLNRLDPLRGF	
fz/Dros	AGLKWGHEAIEENKSHLFLVAVVPAVQITISVLALAKVEGDILSGVCVGLDTHSLGAF	
fz2/Dros/	AGLKWGNEAIIKHSQYFHLAAWLIPVQSVAVLLLSAVDGDPIIGICYVGNLNPDLKTF	
	: ** : : ** * : : *	

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Figure 2c

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                                TM5
fz3/mouse  VLAPLCLYVVVGVSLLLAGIISLNRVRIEIPLEKE-----NQDKLVKFMIRIGVFSILYL
fz4/mouse  VVAPLFTYLVIGTLFIAAGLVALFKIRSNLQK-DG----TKTDKLERLMVKIGVFSVLYT
fz8/mouse  VLAPLVIYLFIGTMFLLAGFVSLFRIRSVIKQQGGP---TKTHKLEKLMIRLGLFTVLYT
fz5/human  VLGPLVLYLLVGTFLFLLAGFVSLFRIRSVIKQ-GG----TKTDKLEKLMIRIGIFTLLYT
fzd9/human VLVPLSGYLVLGSSFLLTG FVALFHIRKIMKT-GG----TNTEKLEKLMVKIGVFSILYT
fz1/rat    VLAPLFVYLFIGTSFLLAGFVSLFRIRTIMKH-DG----TKTEKLEKLMVRIGVFSVLYT
fz2/rat    VLAPLFVYLFIGTSFLLAGFVSLFRIRTIMKH-DG----TKTEPLERLMVRIGVFSVLYT
fz/Dros    LILPLCIYLSIGALFLLAGFISLFRIRTVMKT-DG----KRTDKLERLMRIGFFSGLFI
fz2/Dros/  VLAPLFVYLVIGTTFLMAGFVSLFRIRSVIKQQGGVGAGVKADKLEKLMIRIGIFSILYT
          : *      : : * : . : * : : : . : : : : : * : * :
          ----- TM6 -----> extracellular domain loop 3 <-----
fz3/mouse  VP LLVVIGCYFYEQAYRGIWETTWIQERCREYHIPCPYQVTQMS-----RPDLILFLM
fz4/mouse  VPATCVIACYFYEISN WALFRYSADDS-----NMAVEML
fz8/mouse  VPA AVVVACLFYEQHNRPWEATHNCPCLRDLPDQARR-----PDYAVFML
fz5/human  VPASIVVACYLYEQHYRESWEAALTCACPGHDTGQPRAK-----PEYWVLMML
fzd9/human VPATCVIVCYVYERLNMDFWRLRATEQPCAAAAGPGGRRDCSLP----GGSVPTVAVFML
fz1/rat    VPATIVIAICYFYEQAFRDQWERSWVAQSCSKSYAIPCPHLQGGGGVPPHPMSPDFTVMI
fz2/rat    VPATIVIAICYFYEQAFREHWERSWVSQHCKSLAIPCPAHYT-----PRTSPDFTVYMI
fz/Dros    LPAVGLLGCLFYEYNEDEWMIQWHRDICKPFSIPCPAARAPGS----PEARPIFQIFMV
fz2/Dros/  VPATIVIGCYLYEAAYFEDWIKALACPCAQVK--GPGKK-----PLYSVLML
          *      : . . *
          ----- TM7 -----
fz3/mouse  KYLMALIVGIPSIFWVGSKKTCFEWASFFHGRRKKEIVNESRQVLQEPDFAQSLLRDPNT
fz4/mouse  KIFMSLLVGITSGMWIWSAKTLHTWQKCS-----NRLVNSGKVK-----REKRG
fz8/mouse  KYFMCLVVGITSGVWVWSGKTLESWRALC-----TRCCWASKGAAGVAGAGGSG
fz5/human  KYFMCLVVGITSGVWVWSGKTIVESWRRFT-----SRCCCRPR-----RGHK-
fzd9/human KIFMSLVVGITSGVWVWSKTFQTWQSLC-----YRKIAAGRARA-----KACRA
fz1/rat    KYLMTLIVGITS GFVIWSGKTLNSWRKFY-----TRLTNSK-----QGETT
fz2/rat    KYLMTLIVGITS GFVIWSGKTLHWRKFY-----TRLTNSR-----HGETT
fz/Dros    KYLC SMLVGVITSSVWLYSSKTMVSWRNFV-----ERLQGKEPRT-----RAQAY
fz2/Dros/  KYFMALAVGITS GFVIWSGKTLESWRRFW-----RRLGAPDRTGANQALIKQR
          : : : * : . : * * * :

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Figure 3

Sequence alignment of a portion of the aminoterminal extracellular region of human
Frizzled receptors

HF21 VGQNTSDKGT---PSLLPEFWTSNPQHGGGGHGRG-----GFPGGAG----ASERGK¹FSCPR
HF22 VGQNHSEDGA-----PALLTTAPPPGLQPGAGGTPG-----GPGGGGAPP²RYATLEHPFHC
HF23 LVDLNLAG-----EPTEGAPV-----AVQRDYG-----FWC
HF24 CMEGPGD-----EE-----VPLPHKTPI-----QP
HF25 CMDYNRSEATTAPPRPFP³AKPTLPG-----PPGA-----PASGG-----ECPAGGP⁴PFV-----CKC
HF26 TFDPHTEF-----LGPQKKTE-----QVQRDIG-----FWC
HF27 VGQNTSDGSGGPGGGPTAYPTAPYLPDL⁵PFTALPPG-----ASDGRGRPAF-----PFSC
HF28 CMDYNRTDLTTAAPSPPRRLLPPPP-GEQPPSGSGHGRPPGARPPH⁶RGGGRGGGGDAAAPPARGGGGGKARPPGGGAAP---CEPGCQC
HF29 CMEAPENA-TAGPAEPHKGLGMLPV-----APRPARPPG-----DLGP
HF30 NYLCMEAPNN---GSDEPTRGSGLFPP-----LFRPQRPHSAQ---EHP

Figure 4

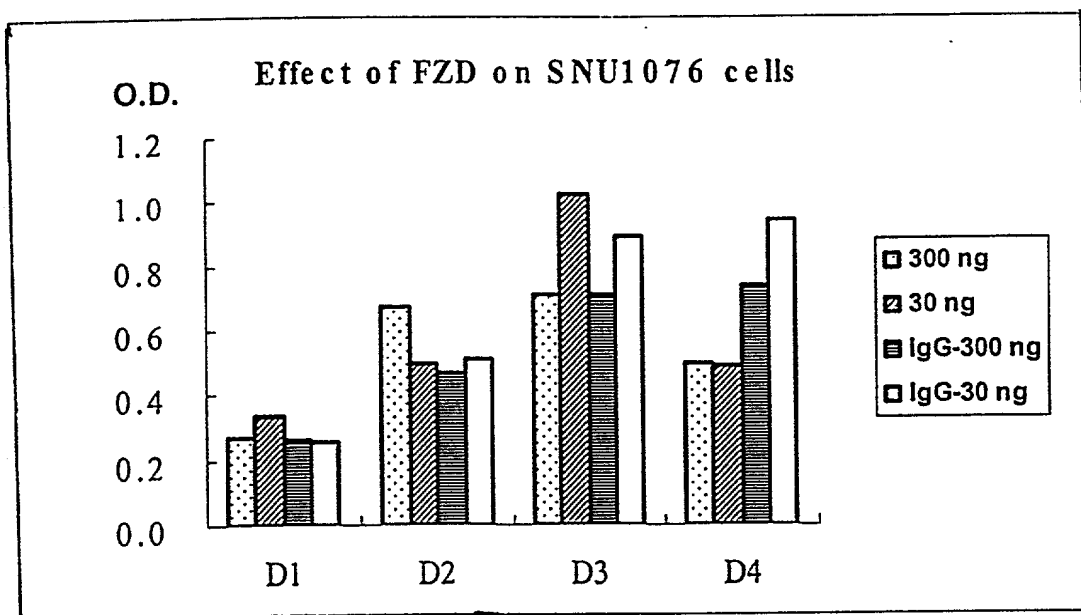


Figure 5

Effect of antibodies SNU 1076 Cells

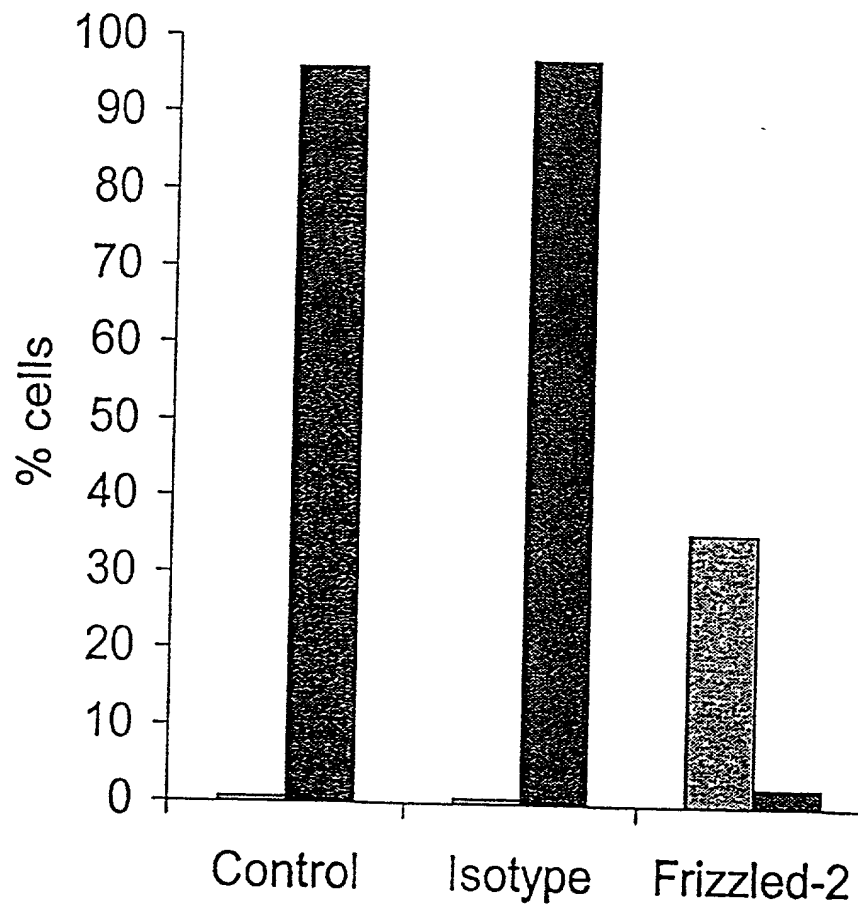
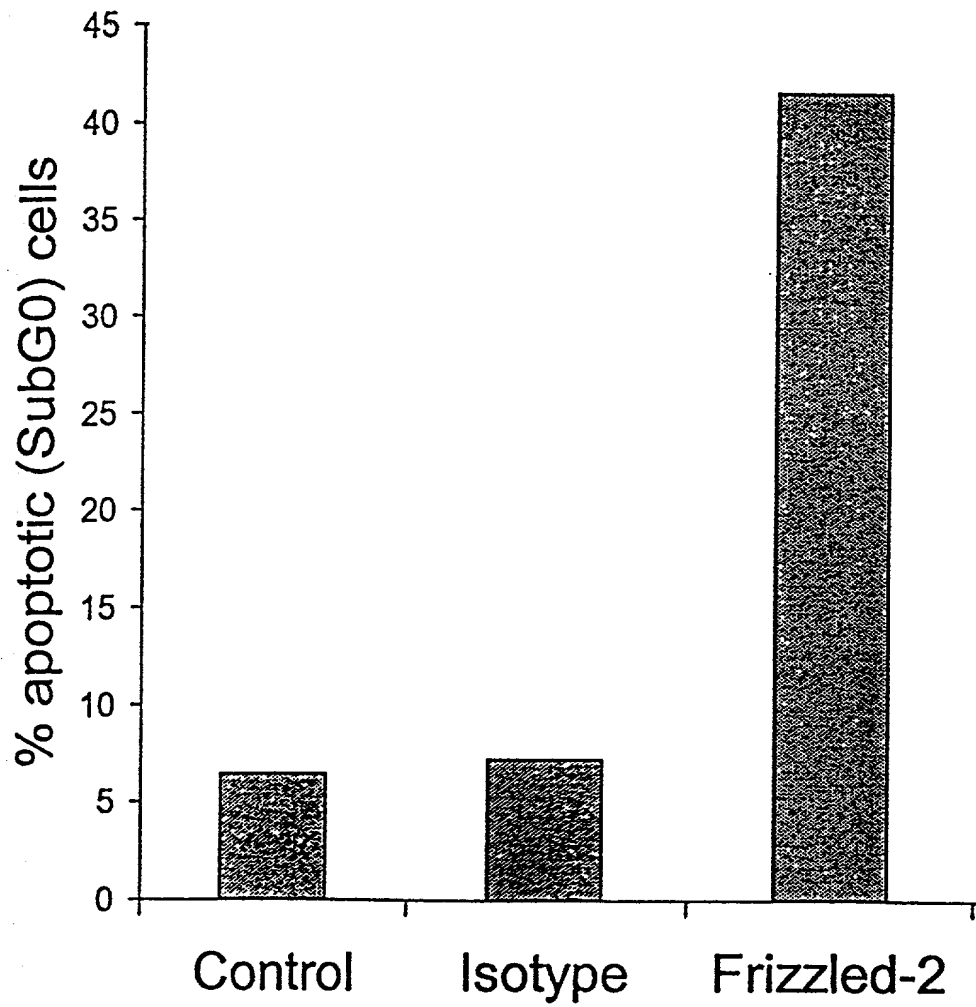


Figure 6

Effect of antibodies on SNU 1076 cells



[illegible]

→ **amino terminal domain**

HFZ1	MAEEEAPKKSRAAGGGASWELCAGALSARLAEEGSGDAGGRRRPPVDPRLRARQLLLLLLW
MFZ1	MAEEEAAPSESRAAGR-LSLELCAEALPGRREEVGHEDTASHRRPRADPRRWASGLLLLLLW
HFZ2	-----MRPRSALPRLLPLL
HFZ3	-----MAMTWIVFSLWPLTV
MFZ3	-----MAVSWIVFDLWLLTV
HFZ4	-----MAWRGAGPSVPGAPGGVGLSLGLLLQ
MFZ4	-----MAWPGTGPSSRGAPGGVGLRLGLLLQ
HFZ5	-----MARPDPSAPPSLL--LLL
HFZ6	-----MEMFTFLTCT
MFZ6	-----MERSPFLLACI
HFZ7	-----MRDPGAAAPLSSLGLCALVLA
MFZ7	-----MRGPGTAASHSPGLCALVLA
HFZ8	-----MEWGYLLEVTSLLAALAL
MFZ8	-----MEWGYLLEVTSLLAALAV
HFZ9	-----MAVAPL-RGALLLWQLLA
MFZ9	-----MAVPPLLRGALLLWQLLA
HFZ10	-----MQRPGPRLWLVLQ

HFZ1	LLEAPL	LLGVRAQAAGQGPGQGP	PGQPPPPPPQ	QQQSGQQYNGERG	ISVPDHGYCQ	PIS
MFZ1	LLEAPL	LLGVRAQAAGQVSG---	PGQQAPPPPPQ	PQQSGQQYNGERG	ISIPDHGYCQ	PIS
HFZ2	LLPA---	-----A-----	-----	GPAQFHGEKGI	SIPDHGFCQ	PIS
HFZ3	FMGHI-----	-----	-----	GGHSLFS-----	-----	CEPIT
MFZ3	FLGQI-----	-----	-----	GGHSLFS-----	-----	CEPIT
HFZ4	LLLLLG-----	-----	-----	PARGFGDEEE-----	-----	RRCDPIR
MFZ4	FLLLLR-----	-----	-----	PTLGFGDEEE-----	-----	RRCDPIR
HFZ5	LAQLVG-----	-----	-----	RAAAASKAPV-----	-----	CQEIT
HFZ6	FLPLL-----	-----	-----	RGHSLFT-----	-----	CEPIT
MFZ6	LLPLV-----	-----	-----	RGHSLFT-----	-----	CEPIT
HFZ7	LLGAL-----	-----	-----	SAGAGAQPYPHGEKGI	SVPDHGFCQ	PIS
MFZ7	LLGAL-----	-----	-----	PTDTRAQPYPHGEKGI	SVPDHGFCQ	PIS
HFZ8	LQRSSG-----	-----	-----	AAAASAKELA-----	-----	CQEIT
MFZ8	LQRSSG-----	-----	-----	AAAASAKELA-----	-----	CQEIT
HFZ9	AGGAAL-----	-----	-----	EIGRFDPERGR--	GAAPCQAVE	
MFZ9	TGGAAL-----	-----	-----	EIGRFDPERGR--	GPAPCQAME	
HFZ10	VMGSCA-----	-----	-----	AISSMDMERP---	GDGKCQPIE	

* 3 5

HFZ1	IPLCTDIAYNQTIMPNLLGHTNQEDAGLEVHQFYPLVKVQCSAELKFFLCSMYAPVCT-V
MFZ1	IPLCTDMAYNQTIMPNLLGHTNQEDAGLEVHQFYPLVKVQCSAELKFFLCSMYAPVCT-V
HFZ2	IPLCTDIAYNQTIMPNLLGHTNQEDAGLEVHQFYPLVKVQCSPELRFFLCSMYAPVCT-V
HFZ3	LRMCQDLPYNTTTFMPNLLNHYDQQTAALAMEPFHPMVNLDCSRDRFRPFLCALYAPICM-E
MFZ3	LRMCQDLPYNTTTFMPNLLNHYDQQTAALAMEPFHPMVNLDCSRDRFRPFLCALYAPICM-E
HFZ4	ISMCQNLGYNVTKMPNVLVGHELQTDABLQLTTFTPLIQYGCSSQLQFFLCSVYVPMCTEK
MFZ4	IAMCQNLGYNVTKMPNVLVGHELQTDABLQLTTFTPLIQYGCSSQLQFFLCSVYVPMCTEK
HFZ5	VPMCRGIGYNLTHMPNQFNHDTQDEAGLEVHQFWPLVEIQCSPDRLRFFLCTMYTPICLPD
HFZ6	VPRCMKMAYNMTFFPNLMGHYDQSIAAVEMEHFLPLANLECSPNIETFLCKAFVPTCI-E
MFZ6	VPRCMKMTYNMTFFPNLMGHYDQGIAAVEMGHFLHLANLECSPNIEMFLCQAFIPTCT-E
HFZ7	IPLCTDIAYNQITILPNLLGHTNQEDAGLEVHQFYPLVKVQCSPELRFFLCSMYAPVCT-V
MFZ7	IPLCTDIAYNQITILPNLLGHTNQEDAGLEVHQFYPLVKVQCSPELRFFLCSMYAPVCT-V
HFZ8	VPLCKGIGYNYTYMPNQFNHDTQDEAGLEVHQFWPLVEIQCSPDRLKFFLCSMYTPICLED
MFZ8	VPLCKGIGYNYTYMPNQFNHDTQDEAGLEVHQFWPLVEIQCSPDRLKFFLCSMYTPICLED

HFZ9 IPMCRGIGYNLTRMPNLLGHTSQGEAAAEALAEFAPLVQYGCHSHLRFLLCSLYAPMCTDQ
 MFZ9 IPMCRGIGYNLTRMPNLLGHTSQGEAAQLAEFSPLVQYGCHSHLRFLLCSLYAPMCTDQ
 HFZ10 IPMCKDIGYNMTRMPNLMGHENQREAAIQLHEFAPLVEYGCHGLRFFLLCSLYAPMCTEQ
 : * : ** * : ** . * * * : * : : * . . . *** : * *

HFZ1 LEQALPPCRSLCERARQGCEALMNKFGFQWPDTLKCEKFPVHG--AGELCVGQNTSDKGT
 MFZ1 LEQALPPCRSLCERARQGCEALMNKFGFQWPDTLKCEKFPVHG--AGELCVGQNTSDKGT
 HFZ2 LEQAIPPCRSICERARQGCEALMNKFGFQWPERLRCEHFPRHG--AEQICVGQNHSEDEGA
 HFZ3 YGRVTLPCRRLCQRAYSECSKLMEMFGVPWPEDMECSRFPDCE--EPYPRLVDLNLVAG---
 MFZ3 YGRVTLPCRRLCQRAYSECSKLMEMFGVPWPEDMECSRFPDCE--EPYPRLVDLNLVAG---
 HFZ4 INIPIGPCGGMCLSVKRRCEPVLKEFGFAWPESLNCSEKFPQON--DHNHMCMEGPGD----
 MFZ4 INIPIGPCGGMCLSVKRRCEPVLKEFGFAWPESLNCSEKFPQON--DHNHMCMEGPGD----
 HFZ5 YHKPLPPCRSVCERAKAGCSPLMRQYGFAPWPERMSCDRLPVLGRDAEVL CMDYNRSEATT
 HFZ6 QIHVVPPCRKLCEKVYSDCKKLIDTFGIRWPEELECRLQYCD--ETVPVTFDPHTEF---
 MFZ6 QIHVVLPCKRLCEKIVSDCKKLMDTFGIRWPEELECRLPHCD--DTVPVTSHPHTEL---
 HFZ7 LDQAIPPCRSICERARQGCEALMNKFGFQWPERLRCEHFPRHG--AGEICVGQNTSDGSG
 MFZ7 LDQAIPPCRSICERARQGCEALMNKFGFQWPERLRCEHFPRHG--AGEICVGQNTSDGSG
 HFZ8 YKKPLPPCRSVCERAKAGCAPLMRQYGFAPWDRMRCDRLPEQG--NPDTLCMDYNRDTLTT
 MFZ8 YKKPLPPCRSVCERAKAGCAPLMRQYGFAPWDRMRCDRLPEQG--NPDTLCMDYNRDTLTT
 HFZ9 VSTPIACRPMCEQARLRCAPIEQFNFGWPDSDLCARLPTRN--DPHALCMEAPENA--TA
 MFZ9 VSTPIACRPMCEQARLRCAPIEQFNFGWPDSDLCARLPTRN--DPHALCMEAPENA--TA
 HFZ10 VSTPIACRVMCEQARLKCSPIEQFNFKWPDSDLCRKLPNKN--DPNYLCMEAPNN----
 . * : * * : : . . . ** : : * . . .

HFZ1 PT---PSLLPEFWTSNPQHGGGGHGRG-----
 MFZ1 PT---PSLLPEFWTSNGQHGGGGYRG-----
 HFZ2 -----PALLTTAPPPGLQPGAGGTPG-----
 HFZ3 ---EPTGAPV-----A
 MFZ3 ---DPTGAPV-----A
 HFZ4 -----EE-----V
 MFZ4 -----EE-----V
 HFZ5 APPRPFPAKPTLPG-----PPGA-----PASGG-----
 HFZ6 ---LGPQKKTE-----Q
 MFZ6 ---SGPQKKSD-----Q
 HFZ7 GPGGGPTAYPTAPYLPDLPTALPPG-----
 MFZ7 GAGGSPTAYPTAPYLPDPPTAMSP-----
 HFZ8 AAPSPRRRLPPPPP--GEQPPSGSGHGRPPGARPPHRRGGGGGGGDAAPARGGGGGGK
 MFZ8 AAPSPRRRLPPPPPPGEQPPSGSGHSRPPGARPPHRRGGSSRGSGDAAAAPPSRGG----K
 HFZ9 GPAEPHKGLGMLPV-----A
 MFZ9 GPTEPHKGLGMLPV-----A
 HFZ10 GSDEPTRGSGFLFP-----L

HFZ1 GFPGGAG---ASERKGFSCPRALKVPSYLYNHYFLGEKDCGAPCEPTKVYGLMYFGPEEL
 MFZ1 GYPGGAG---TVERGKFSCPRALRVPSYLYNHYFLGEKDCGAPCEPTKVYGLMYFGPEEL
 HFZ2 GPGGGGAPPRYATLEHPFHCPRVLKVPSYLSYKFLGERDCAAPCEPARPDGSMFFSQEET
 HFZ3 VQRDYG-----FWCPRELKIDPDLGYSFLHVRDCSPPCP---NMYFR--REEL
 MFZ3 VQRDYG-----FWCPRELKIDPDLGYSFLHVRDCSPPCP---NMYFR--REEL
 HFZ4 PLPHKTPI-----QPGECHSVGTNSDQYIWKRLNLCVLCGYDAGLY--SRSAK
 MFZ4 PLPHKTPI-----QPGECHSVGSNSDQYIWKRLNLCVLCGYDAGLY--SRSAK
 HFZ5 ECPAGGPFV-----CKCREPFVPILKESHPLYNKVRTGQVPNCAPVPCYQPSFSADER
 HFZ6 VQRDIG-----FWCPRHLKTSGGQGYKFLGIDQCAPP CP---NMYFK--SDEL
 MFZ6 VPRDIG-----FWCPKHLRTSGDQGYRFLGIEQCAPP CP---NMYFK--SDEL
 HFZ7 ASDGRGRPAF-----PFSCPRQLKVPPYLGYRFLGERDCGAPCEPGRANGLMYFKEER
 MFZ7 -SDGRGRLSF-----PFSCPRQLKVPPYLGYRFLGERDCGAPCEPGRANGLMYFKEER
 HFZ8 ARPPGGGAAP---CEPGCQCRAPMVSVSSERHPLYNRVKTGQIANCALPCHNPFSSQDER
 MFZ8 ARPPGGGAAP---CEPGCQCRAPMVSVSSERHPLYNRVKTGQIANCALPCHNPFSSQDER
 HFZ9 PRPARPPG-----DLGPGAGGSGTCENPEKFQYVEKSRSCAPRCGPGVEVFWSRDDK
 MFZ9 PRPARPPG-----DSAPGPGSGGTCENPEKFQYVEKSRSCAPRCGPGVEVFWSRDDK
 HFZ10 FRPQRPHSAQ---EHPLKDGGPGRGCDNPGKFHHVEKSASCAPLCTPGVDVYWSREDK

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HFZ1 RFSRTWIGIWSVLCCASTLFTVLTYLVDMMRRFSYPERPIIFLSGCTAVAVAYIAGFLLLE
 MFZ1 RFSRTWIGIWSVLCCASTLFTVLTYLVDMPRFSYPERPIISLSGCTAVAVAYIAGFLLLE
 HFZ2 RFARLWILTWSVLCCASTFTFTVTTYLVDMMQRFYPERPIIFLSGCTMVSVAYIAGFVLQ
 HFZ3 SFARYFIGLISIIICLSATLFTFLTFLIDVTRFRYPERPIIFYAVCYMMVSLIFFIGFLLLE
 MFZ3 SFARYFIGLISIIICLSATLFTFLTFLIDVTRFRYPERPIIFYAVCYMMVSLIFFIGFLLLE
 HFZ4 EFTDIWMAVWASLCFISTTFTVLTLFLIDSSRFSYPERPIIFLSMCYNIYSIAYIVRLTVG
 MFZ4 EFTDIWMAVWASLCFISTTFTVLTLFLIDSSRFSYPERPIIFLSMCYNIYSIAYIVRLTVG
 HFZ5 TFATFWIGLWSVLCFISTTFTVATFLIDMDTFRYPERPIIFLSACYLVCVSLGFLVRLVVG
 HFZ6 EFAKSFIGTVSIFCLCATLFTFLTFLIDVRRFRYPERPIIYYSVCSIVSLMYFIGFLLG
 MFZ6 DFAKSFIGTVSIFCLCATLFTFLTFLIDVRRFRYPERPIIYYSVCSIVSLMYFVGFLLG
 HFZ7 RFARLVWGVWSVLCCASTLFTVLTYLVDMMRRFSYPERPIIFLSGCTFMVAVAHVAGFLLLE
 MFZ7 RFARLVWGVWSVLSCASTLFTVLTYLVDMMRRFSYPERPIIFLSGCTFMVAVAHVAGFLLLE
 HFZ8 AFTVFWIGLWSVLCFVSTFATVSTFLIDMERFKYPERPIIFLSACYLFSVGVYLRLVAG
 MFZ8 AFTVFWIGLWSVLCFVSTFATVSTFLIDMERFKYPERPIIFLSACYLFSVGVYLRLVAG
 HFZ9 DFALVWMAVWSALCFSTTFTVLTLFLLEPHRFQYPERPIIFLSMCYNVYSLAFLIRAVAG
 MFZ9 DFALVWMAVWSALCFSTTFTVLTLFLLEPHRFQYPERPIIFLSMCYNVYSLAFLIRAVAG
 HFZ10 RFAVVWLAIWAVLCFFSSAFTVLTLFLIDPARFRYPERPIIFLSMCYCVSVGYLIRLFAAG

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↳ extracellular domain loop 1

HFZ1 DRVVCNDK-----FAEDGARTVAQGTCK
 MFZ1 DRVVCNDK-----FAEDGARTVAQGTCK
 HFZ2 ERVVCNER-----FSEDGYRTVVQGTCK
 HFZ3 DRVACNAS-----I---PAQYKASTVTQGSKN
 MFZ3 DRVACNAS-----S---PAQYKASTVTQGSKN
 HFZ4 RERISCDF-----EEAEPVLIQEGLEK
 MFZ4 RERISCDF-----EEAEPVLIQEGLEK
 HFZ5 HASVACS-----RE-----HNHIHYETTGP
 HFZ6 DSTACNKA-----D---EKLELGDVTVLGSKN
 MFZ6 NSTACNKA-----D---EKLELGDVTVLGSKN
 HFZ7 DRAVCVER-----FSDDGRTVAQGTCK
 MFZ7 DRAVCVER-----FSDDGRTVAQGTCK
 HFZ8 HEKVACSGGAPGAGGAGGAGGAAA-GAGAAGAGAGGPGGRGEYELGAVEQHVRVYETTGP
 MFZ8 HEKVACSGGAPGAGGRGGAGGAAAAGAGAARGASSPGARGEYELGAVEQHVRVYETTGP
 HFZ9 AQSACD-----QEAGALYVIQEGLEN
 MFZ9 AQSACD-----QEAGALYVIQEGLEN
 HFZ10 AESIACD-----RDSGQLYVIQEGLES

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HFZ1 EGCTILFMMLYFFSMASSIWWVILSLTWFLAAGMKWGHEAIEANSQYFHLAAWAVPAIKT
 MFZ1 EGCTILFMMLYFFSMASSIWWVILSLTWFLAAGMKWGHEAIEANSQYFHLAAWAVPAIKT
 HFZ2 EGCTILFMMLYFFSMASSIWWVILSLTWFLAAGMKWGHEAIEANSQYFHLAAWAVPAVKT
 HFZ3 KACTMLFMILYFFTMAGSVWWVILTITWFLAAVPKWGSEAEIKKALLFHASAWGIPGTLT
 MFZ3 KACTMLFMVLYFFTMAGSVWWVILTITWFLAAVPKWGSEAEIKKALLFHASAWGIPGTLT
 HFZ4 TGCAIIFLLMYFFGMASSIWWVILTLTWFLAAGLKWGHEAIEMHSSYFHIAAWAIPAVKT
 MFZ4 TGCAIIFLLMYFFGMASSIWWVILTLTWFLAAGLKWGHEAIEMHSSYFHIAAWAIPAVKT
 HFZ5 ALCTIVFLLVYFFGMASSIWWVILSLTWFLAAGMKWGNEAIAAGYQYFHLAAWLIPSVKS
 HFZ6 KACTVLFMLLYFFTMAGTVWWVILTITWFLAAGRKWSCEAIEQKAVWFHAWGTPGFLT
 MFZ6 KACSVMFLYFFTMAGTVWWVILTITWFLAAGRKWSCEAIEQKAVWFHAWGAPGFLT
 HFZ7 EGCTILFMVLYFFGMASSIWWVILSLTWFLAAGMKWGHEAIEANSQYFHLAAWAVPAVKT
 MFZ7 EGCTILFMVLYFFGMASSIWWVILSLTWFLAAGMKWGHEAIEANSQYFHLAAWAVPAVKT
 HFZ8 ALCTVFLVLYFFGMASSIWWVILSLTWFLAAGMKWGNEAIAAGYQYFHLAAWLIPSVKS
 MFZ8 ALCTVFLVLYFFGMASSIWWVILSLTWFLAAGMKWGNEAIAAGYQYFHLAAWLIPSVKS
 HFZ9 TGCTLVFLLLYFFGMASSLWVVLTLTWFLAAGKKWGHEAIEAHGSYFHLAAWGLPALKT
 MFZ9 TGCTLVFLLLYFFGMASSLWVVLTLTWFLAAGKKWGHEAIEAHGSYFHLAAWGLPALKT
 HFZ10 TGCTLVFLVLYFFGMASSLWVVLTLTWFLAAGKKWGHEAIEANSYFHLAAWIPAVKT

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↳ extracellular domain loop 2 ←

HFZ1 ITILALGQVDGDLVSGVCFVGLNNVDALRGFVLAPLFVYLFIGTSFLLAGFVSLFRITI
 MFZ1 ITILALGQVDGDLVSGVCFVGLNNVDALRGFVLAPLFVYLFIGTSFLLAGFVSLFRITI
 HFZ2 ITILAMGQIDGDLVSGVCFVGLNSLDPLRGFVLAPLFVYLFIGTSFLLAGFVSLFRITI
 HFZ3 IILLAMNKIEGDNISGVCVGLYDVALRYFVLAPLCYVVGVSLLLAGIISLNRVRIE
 MFZ3 IILLAMNKIEGDNISGVCVGLYDVALRYFVLAPLCYVVGVSLLLAGIISLNRVRIE

Figure 8d

|→ extracellular domain loop 2 ←|

HFZ4 IVILIMRLVDADELTLGLCYVGNQNLDAITGTFVAPLFTYLVIGTLFIAAGLVALFKIRSN
 MFZ4 IVILIMRLVDADELTLGLCYVGNQNLDAITGTFVAPLFTYLVIGTLFIAAGLVALFKIRSN
 HFZ5 ITALALSSVDGDPVAGICYVGNQNLNSLRFRVGLPLVLYLLVGTLLFLLAGFVSLFRIRSV
 HFZ6 VMLLAMNKVEGDNISGVCFVGLYDLASRYFVLLPLCLCVFVGLSLLLAGIISLNHVRQV
 MFZ6 VMLLAMNKVEGDNISGVCFVGLYDLASRYFVLLPLCLCVFVGLSLLLAGIISLNHVRQV
 HFZ7 ITILAMGQVDGDLSSGVCYVGLSSVDALRGFVLAPLFVYLFIGTSFLLAGFVSLFRIRTI
 MFZ7 ITILAMGQVDGDLSSGVCYVGLSSVDALRGFVLAPLFVYLFIGTSFLLAGFVSLFRIRTI
 HFZ8 IAVLALSSVDGDPVAGICYVGNQSLDNLRGFVLAPLVIYLFIGTMFLLAGFVSLFRIRSV
 MFZ8 IAVLALSSVDGDPVAGICYVGNQSLDNLRGFVLAPLVIYLFIGTMFLLAGFVSLFRIRSV
 HFZ9 IVILTLRKVAGDELTLGLCYVASTDAAALTGFVLVPLSGYLVGLSSFLLTGFVALFHIRKI
 MFZ9 IVVLTLRKVAGDELTLGLCYVASMDDPAALTGFVLVPLSCYLVGLTSFLLTGVALFHIRKI
 HFZ10 ILILVMRRVAGDELTVGCYVGSMDVNALTGFVLIPLACYLVIGTSFILSGFVALFHIRRV

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|→ extracellular domain loop 3

HFZ1 MKH--DGTKTEKLEKLMVRIGVFSVLYTVPATIVIACYFYEQAFRDQWERSWVAQSCKSY
 MFZ1 MKH--DGTKTEKLEKLMVRIGVFSVLYTVPATIVIACYFYEQAFRDQWERSWVAQSCKSY
 HFZ2 MKH--DGTKTEKLERLMVRIGVFSVLYTVPATIVIACYFYEQAFREHWERSWVSQHCKSL
 HFZ3 IPL--EKENQDKLVKFMIRIGVFSILYLVPLLVLVIGCYFYEQAYRGIWETTWIQREREY
 MFZ3 IPL--EKENQDKLVKFMIRIGVFSILYLVPLLVLVIGCYFYEQAYRGIWETTWIQREREY
 HFZ4 LQK--DGTKTDKLERLMVKIGVFSVLYTVPATCVIACYFYEISNWALFRYSADDSNMAV-
 MFZ4 LQK--DGTKTDKLERLMVKIGVFSVLYTVPATCVIACYFYEISNWALFRYSADDSNMAV-
 HFZ5 IKQ--GGTKTDKLEKLMIRIGIFTLLYTVPASIVVACLYEQHYRESWEAALTACPGHD
 HFZ6 IQH--DGRNQEKLLKFMIRIGVFSGLYLVPLVTLGLCYVYELVNRIWETWVSDHCRQY
 MFZ6 IQH--DGRNQEKLLKFMIRIGVFSGLYLVPLVTLGLCYVYELVNRIWETWVSDHCRQY
 HFZ7 MKH--DGTKTEKLEKLMVRIGVFSVLYTVPATIVLACYFYEQAFREHWERTWLLQTCKSY
 MFZ7 MKH--DGTKTEKLEKLMVRIGVFSVLYTVPATIVLACYFYEQAFREHWERTWLLQTCKSY
 HFZ8 IKQDGPPTKTHKLEKLMIRLGLFTVLYTVPAAVVVACLFYEQHNRPRWEATHNCPCLRDL
 MFZ8 IKQDGPPTKTHKLEKLMIRLGLFTVLYTVPAAVVVACLFYEQHNRPRWEATHNCPCLRDL
 HFZ9 MKT--GGTNTKLEKLMVKIGVFSILYTVPATCVIVCYVYERLNMDFWRLRATEQPCTAA
 MFZ9 MKT--GGTNTKLEKLMVKIGVFSILYTVPATCVIVCYVYERLNMDFWRLRATEQPCTAA
 HFZ10 MKT--GGENTDKLEKLMVRIGLFSVLYTVPATCVIACYFXEHLNMDYWKILAAQHKCKM-

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HFZ1 AIPCPHLQAGGGAPPHPPMSPDFTVFMIKYLMTLIVGITSGFWIWSGKTLNSWRKFYTRL
 MFZ1 AIPCPHLQAGGGVPPHPPMSPDFTVFMIKYLMT-----LNSWRKFYTRL
 HFZ2 AIPCP-----AHYTPR--MSPDFTVFMIKYLMTLIVGITSGFWIWSGKTLNSWRKFYTRL
 HFZ3 HIPCP-----YQVTQMSRPDLILFLMKYLMALIVGIPSVFVWVGSKKTCFEWASFFHGR
 MFZ3 HIPCP-----YQVTQMSRPDLILFLMKYLMALIVGIPSVFVWVGSKKTCFEWASFFHGR
 HFZ4 -----EMLKIFMSLLVGITSGMWIWSAKTLHTWQ--KCSNR
 MFZ4 -----EMLKIFMSLLVGITSGMWIWSAKTLHTWQ--KCSNR
 HFZ5 TGQPR----AK-----PEYWVLMKYFMCLVVGITSGVWVWSGKTVESWRRFTSRC
 HFZ6 HIPCP-----YQAKAKARPELALFMIKYLMTLIVGISAVFVWVGSKKTCFEWAGFFKRN
 MFZ6 RIPCP-----YQANPKARPELALFMIKYLMTLIVGISAVFVWVGSKKTCFEWAGFFKRN
 HFZ7 AVPCP----PGHFPPM--SPDFTVFMIKYLMTMIVGITTGFVWWSGKTLQSWRRFYHRL
 MFZ7 AVPCP----PRHFSPM--SPDFTVFMIKYLMTMIVGITTGFVWWSGKTLQSWRRFYHRL
 HFZ8 QPDQA----RR-----PDYAVFMLKYFMCLVVGITSGVWVWSGKTLESWRSLCTRC
 MFZ8 QPDQA----RR-----PDYAVFMLKYFMCLVVGITSGVWVWSGKTLESWRSLCTRC
 HFZ9 AGPGG----RRDCSLPGGSVPTVAVFMLKIFMSLVVGITSGVWVWSGKTFTQWQSLCYRK
 MFZ9 TVPGG----RRDCSLPGGSVPTVAVFMLKIFMSLVVGITSGVWVWSGKTFTQWQSLCYRK
 HFZ10 NNQTK----TLDC--LMAASIPAVEIFMVKIFMLLVVGITSGMWIWSKTLQSWQVCSRR

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HFZ1 TN--SKQGETTV-----
 MFZ1 TN--SKQGETTV-----
 HFZ2 TN--SRHGETTV-----
 HFZ3 RKKEIVNESRQVLQEP-----DFAQSLLRDPNTPIIRKSRGTSTQGTSTHAS
 MFZ3 RKKEIVNESRQVLQEP-----DFAQSLLRDPNTPIIRKSRGTSTQGTSTHAS
 HFZ4 LVNSGKVKREKRGNGW-----VKPGKGSE-----
 MFZ4 LVNSGKVKREKRGNGW-----VKPGKGNE-----
 HFZ5 CC--RPRRGHKSGGA-----MA--AG-D-----
 HFZ6 RKRDPISESRRVLQESCEFFLKHNKSVKHKKKHYKPSHKLKVISKSMGTSTGATANHGT
 MFZ6 RKRDPISESRRVLQESCEFFLKHNKSVKHKKKHGAPGPHRLKVISKSMGTSTGATTNHGT

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Figure 8e

HFZ7	SH--SSKGETAV-----
MFZ7	SH--SSKGETAV-----
HFZ8	CW-ASKGAAVGGGAGA-----TAAGGGGGPGGGGGGGP
MFZ8	CW-ASKGAAVGGAGAGG-----SGPGGSGP-----GP
HFZ9	IA--AGRARAKACRAP-----GSYGRGTHC-----
MFZ9	MA--AGRARAKACRTP-----GGYGRGTHC-----
HFZ10	LKKKSRRKPASVITSG-----GIYKKAQH-----

HFZ1	-----
MFZ1	-----
HFZ2	-----
HFZ3	STQLAMVDDQRSKAGSIHISKVSSYHGSLHRSRDGRYTPCSYRG--MEERLPHGSMS-RLT
MFZ3	STQLAMVDDQRSKAGSVHISKVSSYHGSLHRSRDGRYTPCSYRG--MEERLPHGSMS-RLT
HFZ4	-----TVV-----
MFZ4	-----TVV-----
HFZ5	-----YPEASAALTGRTGPPGPAATYHKQVSLSHV-----
HFZ6	SAVAITSHDYLGOETLITEIQTSPETSMREVKADGASTPRLREQDCGEPASPAASIS-RLS
MFZ6	SAMAIADHDYLGQETSTEVHTSPEASVKEGRADRANTPSAKDRDCGESAGPSSKLSGNRN
HFZ7	-----
MFZ7	-----
HFZ8	GGGGGPGGGGSLYSDVSTGLTWRSKTAS-SVSYPKOMPLSQV-----
MFZ8	GGGGGHGGGGSLYSDVSTGLTWRSKTAS-SVSYPKOMPLSQV-----
HFZ9	-----H--YKAPTIVLHMTKTDPSLENPTL-----
MFZ9	-----H--YKAPTIVLHMTKTDPSLENPTL-----
HFZ10	-----PQKT-HHGKYEIPAQSPTCV-----

HFZ1	-----
MFZ1	-----
HFZ2	-----
HFZ3	DHSRHSSSHRLNEQSRHSSIRDLNPNMTHITHGTSMNRVIEEDGTTA-----
MFZ3	DHSRHSSSHRLNEQSRHSSIRDLNPNMTHITHGTSMNRVIEEDGTTA-----
HFZ4	-----
MFZ4	-----
HFZ5	-----
HFZ6	GEQVDGKG--QAGSVSESARSEGRISPKSDITDTGLAQSNLQVPSSEPSLKGSTSL
MFZ6	GRESRAGGLKERSNGSEGAPSEGRVSPKSSVPETGLIDCSTSAASSPEPTSLKGSTSLP
HFZ7	-----
MFZ7	-----
HFZ8	-----
MFZ8	-----
HFZ9	-----
MFZ9	-----
HFZ10	-----

HFZ1	-----
MFZ1	-----
HFZ2	-----
HFZ3	-----
MFZ3	-----
HFZ4	-----
MFZ4	-----
HFZ5	-----
HFZ6	VHPVSGVRKEQGGGCHSDT
MFZ6	VHSASRARKEQGAGSHSDA
HFZ7	-----
MFZ7	-----
HFZ8	-----
MFZ8	-----
HFZ9	-----

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